

D. knuse

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/954,773A

DATE: 01/14/2003
TIME: 14:52:26

Input Set : N:\paola\US09954773A.raw
Output Set: N:\CRF4\01142003\I954773A.raw

1 <110> APPLICANT: Southern Illinois University
2 Lightfoot, David A
3 Gibson, Paul T
4 Meksem, Khalid
5 <120> TITLE OF INVENTION: SOYBEAN SUDDEN DEATH SYNDROME RESISTANT SOYBEANS, SOYBEAN
CYST NEMATODE
6 RESISTANT SOYBEANS AND METHODS OF BREEDING AND IDENTIFYING RESISTANT PLANTS
7 <130> FILE REFERENCE: 1268/2/2
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/954,773A
9 <141> CURRENT FILING DATE: 2002-01-22
10 <150> PRIOR APPLICATION NUMBER: 09/007,119
11 <151> PRIOR FILING DATE: 1998-01-14
12 <160> NUMBER OF SEQ ID NOS: 20
13 <170> SOFTWARE: PatentIn version 3.1
15 <210> SEQ ID NO: 1
16 <211> LENGTH: 527
17 <212> TYPE: DNA
18 <213> ORGANISM: Glycine max
19 <220> FEATURE:
20 <221> NAME/KEY: misc_feature
21 <222> LOCATION: (1)..(527)
22 <223> OTHER INFORMATION: n is a, c, g, or t/u
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24 tgttcttagat agttcgaat tcaatcaaatttcccaatttaatttgaataaaaaggattca 60
25 tgaatcagg tgatcaagcg aaaaataaagc attaagcgta gaagagaagc aataacattt 120
26 tttttaaaa taataaaaaga gtaattacat aaaaatatgt tcgattacat taaaccccaa 180
27 caaaggatga attagcttc tcatgaccat gggaaaatc aaacttgatg aacaagaaga 240
28 tgaagaagaa tccttaagga taaacactgc cttagctcaa tggctctt agtattttat 300
29 cttcaaaaaa tcccccaagaa cccctaattt tcagtaagaa gcccattttc aatcagaagc 360
30 ccattttcaa tcaagaagcc cattttcaat cagaagccca ttttcaatca gaagcccat 420
31 ttaatcaga agcccatttt caatcagaag cccatttt aattgtattt ccaaaaacttg 480
32 agattcttga acgtaaattt ttagtaaattt gtaatcacct ctgtaaa 527
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 815
36 <212> TYPE: DNA
37 <213> ORGANISM: Glycine max
38 <220> FEATURE:
39 <221> NAME/KEY: misc_feature
40 <222> LOCATION: (1)..(815)
41 <223> OTHER INFORMATION: n is a, c, g, or t/u
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44 gttggagct ctcccatatg gtcgacctgc aggcggccgc actagtgatt cagaagccca 120
45 aagtaacag caataagtaa tcccttgttt ataagatccc agaacttcca gtttatttaa 180

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46	tgaaaatgca ataacatcg	ctagttcac aagtaatata	caaatcgaa catcacattg	240
47	actacaatat atagtacata	aattaacact aagaaacctc	cttgatttga tattatgc	300
48	ttacctatgt tggccacaa	gaatatactc aaatgactt	gccttgatt aaattatcac	360
W--> 49	gatgtacac aaacaaaagat	gatantttgt cgatcaactg	ttcagcacca agagagccct	420
W--> 50	cccacaatc aactcagggtt	ttcacttttg gtgctgaaa	atgagtggca catgaaaag	480
W--> 51	caagagtcnt ctttgacaaa	tgtgcctgcc ganagttatc	antacttact aacaagataa	540
W--> 52	tgagccaaaa catcatctgg	gnatcaacc ttcatgnctt	tntcaagttt atacctatna	600
W--> 53	ntnactangt cttatatttn	canntggta ttacanttac	nantaagttt agcttnaaga	660
W--> 54	aatncaagtt ttngggactc	catgcctngn cnngnnttcn	natccgtcgg ccagggcggn	720
W--> 55	cngnnncact gntnggnagn	cccanttnch cagancacng	ncccnnntcc attccnggn	780
W--> 56	cntcncttc aangacngcc	gngaaancn ngggt		815
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59	<211> LENGTH: 435			
60	<212> TYPE: DNA			
61	<213> ORGANISM: Glycine max			
62	<220> FEATURE:			
63	<221> NAME/KEY: misc_feature			
64	<222> LOCATION: (1)..(435)			
65	<223> OTHER INFORMATION: n is a, c, g, or t/u			
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68	ccatagcaag caaagtatgg	atttgcatt tttcagagac	gaagaactct caacaaacat	120
69	gtttagtacttcaacttgc	aaaactcaac aaatagattt	tttgaacctt aatataataa	180
70	aattcaacag tcttctttaa	tttattctg ctcttacctt	ctcataggat catatagaat	240
71	ttaaccctac aagctctcaa	aaaacaatcc attattatgc	tccttatcca ataaaacaaa	300
W--> 72	accatagagt gattctaaa	atgaagattt acaaaggcaa	aaagttatgc tggntcaata	360
W--> 73	gctctttat aattntcttc	atcttgccacc ntccngcct	taggnggtct ccattgtcaa	420
W--> 74	tccaaaggtn ntcttc			435
76	<210> SEQ ID NO: 4			
77	<211> LENGTH: 183			
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79	<213> ORGANISM: Glycine max			
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81	<221> NAME/KEY: misc_feature			
82	<222> LOCATION: (1)..(183)			
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84	<400> SEQUENCE: 4			
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86	tcgcctat ggccttacag	ggttgcga attagtgtga	aggttaattcg gtaaatggat	120
W--> 87	aatattgtat tcatttnata	tttnatgtat ttacaagtnc	aaggtnataan ctgatgcctg	180
88	agt			183
90	<210> SEQ ID NO: 5			
91	<211> LENGTH: 499			
92	<212> TYPE: DNA			
93	<213> ORGANISM: Glycine max			
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95	<221> NAME/KEY: misc_feature			
96	<222> LOCATION: (1)..(499)			
97	<223> OTHER INFORMATION: n is a, c, g, or t/u			

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 W--> 100 aagcccagag aatatgtgct agcttggaaag tngtgggagg ggagtgtga aacatttac 120
 101 tggtttatga aggtataaca ccaattatta tggtttttg ttaataaaa tgtgaataat 180
 102 tgtcaatcgt gattgcatta tctctcctt actctgtctc ttcacccctt ttacccttt 240
 103 atttgagagg aagaatccat gtagaaaaaa atgatgataa aattgttaga aaatatacg 300
 104 tcatgttaatt agagattcag attataactt agaagacact attatttca tgtaatacta 360
 105 tccacgggta attatacaata ctgacatatt ttcactcaaa atattctggg ttttcattta 420
 W--> 106 tatacattta aataggagct attanccatt gcaagcttgg gtttggaggc cttccgatgc 480
 W--> 107 ctgttggga ttgngacca 499
 109 <210> SEQ ID NO: 6
 110 <211> LENGTH: 500
 111 <212> TYPE: DNA
 112 <213> ORGANISM: Glycine max
 113 <220> FEATURE:
 114 <221> NAME/KEY: misc_feature
 115 <222> LOCATION: (1)..(500)
 116 <223> OTHER INFORMATION: n is a, c, g, or t/u
 117 <400> SEQUENCE: 6
 W--> 118 nagnccggga acccacacat acagacaatt aaaaccgttg gatgaaaatc atactactca 60
 W--> 119 taaattgaaa atatatacgt aagancttca tctaacagtg ctagtcgaag aatgcgtaaa 120
 W--> 120 tgccagggnat ccattccat actaaaatgg acaaaaactta tattttttt ttagcggcaa 180
 W--> 121 acgttaatta ttaatttttt ttagtacaag ggatcaaacc angaccttcc ccttcttcc 240
 122 atctttcttg accacccaac caaccttata tctccacaaa acttattata tgggtttctt 300
 W--> 123 cggggactat cagaatttggaa gtttaacctc gggcantcaa tctacataat ccttgatttn 360
 W--> 124 atttngtggaa gttctaaagc cacaggcatt atttatntt ttnnttctgn agtaaccnc 420
 W--> 125 catatgttgg tnnataaggg tangnatnaa aatncnttgg ntggtnncna tttgcncttn 480
 W--> 126 cnaggccggg gatggntttt 500
 128 <210> SEQ ID NO: 7
 129 <211> LENGTH: 189
 130 <212> TYPE: DNA
 131 <213> ORGANISM: Glycine max
 132 <220> FEATURE:
 133 <221> NAME/KEY: misc_feature
 134 <222> LOCATION: (1)..(189)
 135 <223> OTHER INFORMATION: n is a, c, g, or t/u
 136 <400> SEQUENCE: 7
 W--> 137 nnacaanana ncaggggatc ctctagagtc gacctgcagt gatactagaa ctnaatgaac 60
 W--> 138 agggagagag agagagagag agananntaa nataacgtg aagctctccc tattgacgg 120
 W--> 139 gttcattgtt gcaatagcat cgttatctct tattattgct gttcatcat natctcaatt 180
 140 ccagtggca 189
 142 <210> SEQ ID NO: 8
 143 <211> LENGTH: 724
 144 <212> TYPE: DNA
 145 <213> ORGANISM: Glycine max
 146 <220> FEATURE:
 147 <221> NAME/KEY: misc_feature
 148 <222> LOCATION: (1)..(724)
 149 <223> OTHER INFORMATION: n is a, c, g, or t

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152 ctaatttatc atatttttg tatattcagg aattaaattt aattttcat ccttcaatac 120
153 taacttatta acgtttcaca ttttcaaaga cgagtctagc tatttataat ttttttcct 180
154 aaaaatattt ttttgcctc ataaatatga aatattaa aattcgttcc taatttttt 240
155 ttc当地aaagcat ctttccctc cacaatttgg aatgtatca ttttttttg tt当地aaagtt 300
156 taaataaattt tgaacctaat atgacattt atatcggtt tacatataac tgatataaac 360
157 atcaagttt ttatataat gatacctata actgatataca aatgtgacaa ttatataat 420
158 aattaatgtt aaaaagtc当地 aatataattt tattttgagt caaaataaa tatattttaa 480
W--> 159 ttatTTTgaa gatgaaaaag gataaatttta aaacatttggt gt当地atgtt aaaaactagat 540
160 gtttttttc ctggttt当地 tgccaaacc当地 atgctattt atttaattt tacctttt 600
W--> 161 ttataatttcc nccacaaaaa aaccgtttgg tggatcaat tt当地antttaaa tt当地nttgg 660
W--> 162 tattaaaaag ananattaat tnggaanggt ct当地ttnaaa acnctncngt cnantaacna 720
163 atct 724
165 <210> SEQ ID NO: 9
166 <211> LENGTH: 801
167 <212> TYPE: DNA
168 <213> ORGANISM: Glycine max
169 <220> FEATURE:
170 <221> NAME/KEY: misc_feature
171 <222> LOCATION: (1)..(801)
172 <223> OTHER INFORMATION: n is a, c, g, or t
173 <400> SEQUENCE: 9
W--> 174 acgcccagtga ntgtataacg actcctatacg ggc当地aattgg ccaagtc当地c cgagctcgaa 60
175 ttc当地tc当地 accgaggatc acgctatacg tatatttata atcaactgtc tcaatagat 120
176 gcacacaccc tatcttcat aaaattacta cacttttaa ttttgc当地 aaaaacacta 180
177 gaaaaactca tt当地aaaca gatgatgtac tttt当地act ct当地c当地ctct 240
178 attatataattt gattt当地attt tattt当地at tattttttg tt当地gtctca tt当地attat 300
179 tttt当地attt ggtatccggc cctctatgt cggccgcatg cataagctt agtattctat 360
180 agtgtcaccc aaatagctt gctatcat ggtcatagct gtttctgtg tgaatttgg 420
W--> 181 atccgctcac aattccacac aacatacgag cc当地aaggcat aaatgtt当地a gcttgggtn 480
W--> 182 ccttaatgatc gagctactc acat当地atgg cctt当地ctc actgccc当地t ttc当地gtcng 540
W--> 183 gaaacctgtc ct当地c当地ctg cat当地atggaa tc当地ccaacc cnccgggana agc当地tttgc 600
W--> 184 ntatggc当地 tttt当地ctc nt当地actcgct gctatc当地t gctatc当地t nt当地c当地nttgc 660
W--> 185 cgc当地acggt atc当地ncac tc当地aangnng taaatacggt tatccatc当地a accnnggg 720
W--> 186 naaccnnggaa aaaaacatgt nanccaaag gcnccaaag gccc当地aaaag ctnn当地aaaag 780
W--> 187 gccc当地nttgc tt当地nttntt n 801
189 <210> SEQ ID NO: 10
190 <211> LENGTH: 809
191 <212> TYPE: DNA
192 <213> ORGANISM: Glycine max
193 <220> FEATURE:
194 <221> NAME/KEY: misc_feature
195 <222> LOCATION: (1)..(809)
196 <223> OTHER INFORMATION: n is a, c, g, or t
197 <400> SEQUENCE: 10
W--> 198 nnnnnnttggt aaacgacgca gt当地atgtat acgaccacta tagggcaatg gccaatc当地gg 60
199 cccgagctcgac attc当地ctc当地 ctc当地ggat ct当地t当地atgt tggatctac ttaatata 120
200 tctt当地tactt ttaat当地tta agtcatatctc ct当地tggact catatataag caaaagatgt 180

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201	gtcttgtatg tcggacttaa atataagcaa atctaactaa ttttgccta tttaatactt	240
202	tcattcctaa aacacccttc atttaattct aattctattt ccaataactc ttttttattc	300
203	atgataacaa gttccaatga aggacattt agaaataacc ttatTTTTtta ttgagatta	360
204	gtaaaattaa atgatgtgaa ctaactttct taattaatgt gaaatttagtt attttttctt	420
205	atatacgagt ccaaaggag taccaaattt cacaatgtt ctaaaatgtt ttatatgctt	480
W--> 206	c tttttaatt c atcttgct g cata n tac t tagt a ct g tgct c t g at c cggcc c t c t c 540	540
207	agatgcggcc gcatgcataa gcttgagttt ctata g tgc cctaaat g tc ttgg c gt t at c 600	600
W--> 208	atgg t cat g atg t tt c ng t gt g aa t tg t tat c cg c tc acaatt c ac acaacata c g 660	660
W--> 209	ancc g ga g c ataaa g at g t taag g cc g ng g tg c t a at g agt g ag g ct a ctc a catt a 720	720
W--> 210	tt g c g tt g cg c tc a ct g ccc g ct t cc n att c gg g aa a ct g t c t tg n c a nc t gc a t t at g 780	780
W--> 211	aat c ng g cc a acc c nc g ng g aaa g gc g 809	809
213 <210>	SEQ ID NO: 11	
214 <211>	LENGTH: 810	
215 <212>	TYPE: DNA	
216 <213>	ORGANISM: Glycine max	
217 <220>	FEATURE:	
218 <221>	NAME/KEY: misc_feature	
219 <222>	LOCATION: (1)..(810)	
220 <223>	OTHER INFORMATION: n is a, c, g, or t/u	
221 <400>	SEQUENCE: 11	
W--> 222	acng g cc a gt g aatt g taata c gact c ct t at a gg g caattt g g c ca a gt g tc g 60	60
223	aatt c gt g ca c c tc g agg g ga t t tataat a t tt t ct g ac g ac tac t tttt t ta tt t ag t tc g 120	120
224	agagg g g g ct g at t tt g aga a a ac a cat c at c cat g tt a taa agt g cc g tt a gatt c ca g ct 180	180
225	at t gt t ca c aca t t cat c ctt acat a t g aga at t at c ct t at aag g ct g aa a c ta a ct t tt t ac 240	240
226	aa a caa a cat c g c acc g ga acc att a aa g ttt g a ct a at a t cc g gg g tata at g ac c tt aa 300	300
227	tt c ag g aa att t ca c ata a ata a act a aa g ta agt g tt t attt tatt t at g tc t g gatt t act 360	360
228	gc a caa a act a a a caa a g t ttt g att a g a cata aaaa a a cca a at g c t g tg g aaaa 420	420
W--> 229	taag g aa at gg t gg t cat a ta g aca a gt t tc t t tt t ct g ttt t c t tt a attt g c a gt c na a g 480	480
W--> 230	ccat c ang g a g t tc at gt aa t t aa cc aa ac t ag ac gt tt ga c tt t gg g ttt t at c tt t tt g 540	540
231	tag a at g ca a g g ca a gt tc at c ata a at c tg g c att g gg g a cag t ta g tt ta a ct ccc gc 600	600
W--> 232	cg cc aa at tt g t t aaa at tt g na a ata a at c ac c ac c tt aa at c at a tt g tc an t c att 660	660
W--> 233	tt g tt tt ang t tat c ta a tt t att t tt t tac tt ta c nt c tt t ata a at nt ca at g at g 720	720
W--> 234	gg ac cc cc aaa a a tt a ta aaa tac nt tna ag c nt tatt tt at tatt a att aa nc ct ta att 780	780
W--> 235	ata a tt aa aaa att c att tt ta att tt tt ta an 810	810
237 <210>	SEQ ID NO: 12	
238 <211>	LENGTH: 777	
239 <212>	TYPE: DNA	
240 <213>	ORGANISM: Glycine max	
241 <220>	FEATURE:	
242 <221>	NAME/KEY: misc_feature	
243 <222>	LOCATION: (1)..(777)	
244 <223>	OTHER INFORMATION: n is a, c, g, or t/u	
245 <400>	SEQUENCE: 12	
W--> 246	an a ng g att g n cag g t at ttt ag gt g ac a ta c tag a aa a ta c t a ca a g c tt t at g c at g cc g cc g 60	60
247	cat c ta g agg g cc g gat c t t c g gt g aa g c aaa a tt g a agt t tt t tc g tc t at t tt t at 120	120
248	caa a tt c ttt a a t g aaa g t ta a tt c ata a a a at t tt t ta g t ga a g ca a tt t tt c ac a g 180	180
249	tt t att t tt t ta aaaa att ac ac g tt t tt c a a ta c aa a at t a ca a at t at t ata g gt ta 240	240
250	ta a tt a at t at tt aaa att c at a ta a aa g a t g act t att a ta a g t tg at a a tg ta att a 300	300
251	tt t ta c act a t t aa a ct c at t t tac g ta t ct t ag g c a ca ac a ta c att t tt t cat g a 360	360

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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 Seq#:2; N Pos. 785,786,793,797,803,808,810,811
 Seq#:3; N Pos. 354,375,391,396,405,430,431,435
 Seq#:4; N Pos. 46,54,137,144,159,165,170
 Seq#:5; N Pos. 92,445,494
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 Seq#:19; N Pos. 11,18,166,174,199,222,246,268,277,281,304,309,317,326,356
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 Seq#:19; N Pos. 670,679,705,708,710,751,762,771,786,793,802,804,806
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 Seq#:20; N Pos. 712,714,716,727,730,739,742,748,753,773,780,787

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/14/2003
PATENT APPLICATION: US/09/954,773A TIME: 14:52:27

Input Set : N:\paola\US09954773A.raw
Output Set: N:\CRF4\01142003\I954773A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5,6

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/954,773ADATE: 01/14/2003
TIME: 14:52:27Input Set : N:\paola\US09954773A.raw
Output Set: N:\CRF4\01142003\I954773A.raw

L:8 M:270 C: Current Application Number differs, Wrong Format
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:360
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:420
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:480
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:540
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:720
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:780
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:300
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:360
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:420
L:85 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:87 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:120
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:60
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:420
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:480
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:180
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:300
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:360
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:420
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:480
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:138 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:60
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:120
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:480
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:600
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:660
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:420
L:182 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:480
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:540
L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:600
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:660
L:186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:720
L:187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:780
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:480
L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:600
L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:660
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:720
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:780
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:420
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:480

VERIFICATION SUMMARY

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Input Set : N:\paola\US09954773A.raw
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L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:600
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:660
L:234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:720